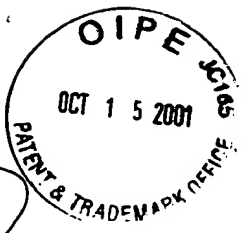


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 Lys Ile Ala Val Tyr Tyr Lys Gly Lys Arg Leu Ala Ser Ala Ala Glu
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 Ser Asn Val His Lys Ala Glu Leu Arg Val Ala Glu Leu Ala Leu Ala
 370 375 380
 Asn Leu Glu Ser Met Ser Phe Ser Lys Met Lys Ala Lys Asn Asn Ser
 385 390 395 400
 Asn Met Arg Arg Arg Leu Glu Gln Asp Thr Ser Asp
 405 410

<210> 4
 <211> 366
 <212> PRT
 <213> *Saccharomyces pombe*

<400> 4

Met Gly Arg Phe Lys Arg His His Glu Gly Asp Ser Asp Ser Ser Ser
 1 5 10 15
 Ser Ala Ser Asp Ser Leu Ser Arg Gly Arg Arg Ser Leu Gly His Lys
 20 25 30
 Arg Ser Ser His Ile Lys Asn Arg Gln Tyr Tyr Ile Leu Glu Lys Lys
 35 40 45
 Ile Arg Lys Leu Met Phe Ala Met Lys Ala Leu Leu Glu Glu Thr Lys
 50 55 60
 His Ser Thr Lys Asp Asp Val Asn Leu Val Ile Pro Gly Ser Thr Trp
 65 70 75 80
 Ser His Ile Glu Gly Val Tyr Glu Met Leu Lys Ser Arg His Asp Arg
 85 90 95
 Gln Asn Glu Pro Val Ile Glu Glu Pro Ser Ser His Pro Lys Asn Gln

100					105					110					
Lys	Asn	Gln	Glu	Asn	Asn	Glu	Pro	Thr	Ser	Glu	Glu	Phe	Glu	Glu	Gly
		115					120					125			
Glu	Tyr	Pro	Pro	Pro	Leu	Pro	Pro	Leu	Arg	Ser	Glu	Lys	Leu	Lys	Glu
	130					135					140				
Gln	Val	Phe	Met	His	Ile	Ser	Arg	Ala	Tyr	Glu	Ile	Tyr	Pro	Asn	Gln
145					150					155					160
Ser	Asn	Pro	Asn	Glu	Leu	Leu	Asp	Ile	His	Asn	Glu	Arg	Leu	Glu	Phe
				165					170					175	
Leu	Gly	Asp	Ser	Phe	Phe	Asn	Leu	Phe	Thr	Thr	Arg	Ile	Ile	Phe	Ser
			180					185					190		
Lys	Phe	Pro	Gln	Met	Asp	Glu	Gly	Ser	Leu	Ser	Lys	Leu	Arg	Ala	Lys
		195					200					205			
Phe	Val	Gly	Asn	Glu	Ser	Ala	Asp	Lys	Phe	Ala	Arg	Leu	Tyr	Gly	Phe
	210					215					220				
Asp	Lys	Thr	Leu	Val	Leu	Ser	Tyr	Ser	Ala	Glu	Lys	Asp	Gln	Leu	Arg
225				230						235					240
Lys	Ser	Gln	Lys	Val	Ile	Ala	Asp	Thr	Phe	Glu	Ala	Tyr	Leu	Gly	Ala
				245					250					255	
Leu	Ile	Leu	Asp	Gly	Gln	Glu	Glu	Thr	Ala	Phe	Gln	Trp	Val	Ser	Arg
			260					265					270		
Leu	Leu	Gln	Pro	Lys	Ile	Ala	Asn	Ile	Thr	Val	Gln	Arg	Pro	Ile	Asp
		275					280					285			
Lys	Leu	Ala	Lys	Ser	Lys	Leu	Phe	His	Lys	Tyr	Ser	Thr	Leu	Gly	His
		290				295					300				
Ile	Glu	Tyr	Arg	Trp	Pro	Ala	Cys	Val	Asp	Gly	Ala	Gly	Gly	Ser	Ala
305					310					315					320
Glu	Gly	Tyr	Val	Ile	Ala	Cys	Ile	Phe	Asn	Gly	Lys	Glu	Val	Ala	Arg
				325					330					335	
Ala	Trp	Gly	Ala	Asn	Gln	Lys	Asp	Ala	Gly	Ser	Arg	Ala	Ala	Met	Gln
			340					345					350		
Ala	Leu	Glu	Val	Leu	Ala	Lys	Asp	Tyr	Ser	Lys	Phe	Ala	Arg		
		355					360					365			

<210> 5
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 <212> PRT
 <213> *Saccharomyces.cerevisiae*

 <400> 5

Met	Gly	Ser	Lys	Val	Ala	Gly	Lys	Lys	Lys	Thr	Gln	Asn	Asp	Asn	Lys	1	5	10	15
Leu	Asp	Asn	Glu	Asn	Gly	Ser	Gln	Gln	Arg	Glu	Asn	Ile	Asn	Thr	Lys	20	25	30	
Thr	Leu	Leu	Lys	Gly	Asn	Leu	Lys	Ile	Ser	Asn	Tyr	Lys	Tyr	Leu	Glu	35	40	45	
Val	Ile	Gln	Leu	Glu	His	Ala	Val	Thr	Lys	Leu	Val	Glu	Ser	Tyr	Asn	50	55	60	
Lys	Ile	Ile	Glu	Leu	Ser	Pro	Asn	Leu	Val	Ala	Tyr	Asn	Glu	Ala	Val	65	70	75	80
Asn	Asn	Gln	Asp	Arg	Val	Pro	Val	Gln	Ile	Leu	Pro	Ser	Leu	Ser	Arg	85	90	95	
Tyr	Gln	Leu	Lys	Leu	Ala	Ala	Glu	Leu	Lys	Thr	Leu	His	Asp	Leu	Lys	100	105	110	
Lys	Asp	Ala	Ile	Leu	Thr	Glu	Ile	Thr	Asp	Tyr	Glu	Asn	Glu	Phe	Asp	115	120	125	
Thr	Glu	Gln	Lys	Gln	Pro	Ile	Leu	Gln	Glu	Ile	Ser	Lys	Ala	Asp	Met	130	135	140	
Glu	Lys	Leu	Glu	Lys	Leu	Glu	Gln	Val	Lys	Arg	Glu	Lys	Arg	Glu	Lys	145	150	155	160
Ile	Asp	Val	Asn	Val	Tyr	Glu	Asn	Leu	Asn	Glu	Lys	Glu	Asp	Glu	Glu	165	170	175	
Glu	Asp	Glu	Gly	Glu	Asp	Ser	Tyr	Asp	Pro	Thr	Lys	Ala	Gly	Asp	Ile	180	185	190	
Val	Lys	Ala	Thr	Lys	Trp	Pro	Pro	Lys	Leu	Pro	Glu	Ile	Gln	Asp	Leu	195	200	205	
Ala	Ile	Arg	Ala	Arg	Val	Phe	Ile	His	Lys	Ser	Thr	Ile	Lys	Asp	Lys	210	215	220	
Val	Tyr	Leu	Ser	Gly	Ser	Glu	Met	Ile	Asn	Ala	His	Asn	Glu	Arg	Leu	225	230	235	240
Glu	Phe	Leu	Gly	Asp	Ser	Ile	Leu	Asn	Ser	Val	Met	Thr	Leu	Ile	Ile	245	250	255	
Tyr	Asn	Lys	Phe	Pro	Asp	Tyr	Ser	Glu	Gly	Gln	Leu	Ser	Thr	Leu	Arg	260	265	270	
Met	Asn	Leu	Val	Ser	Asn	Glu	Gln	Ile	Lys	Gln	Trp	Ser	Ile	Met	Tyr	275	280	285	
Asn	Phe	His	Glu	Lys	Leu	Lys	Thr	Asn	Phe	Asp	Leu	Lys	Asp	Glu	Asn	290	295	300	

Ser Asn Phe Gln Asn Gly Lys Leu Lys Leu Tyr Ala Asp Val Phe Glu
 305 310 315 320
 Ala Tyr Ile Gly Gly Leu Met Glu Asp Asp Pro Arg Asn Asn Leu Pro
 325 330 335
 Lys Ile Arg Lys Trp Leu Arg Lys Leu Ala Lys Pro Val Ile Glu Glu
 340 345 350
 Ala Thr Arg Asn Gln Val Ala Leu Glu Lys Thr Asp Lys Leu Asp Met
 355 360 365
 Asn Ala Lys Arg Gln Leu Tyr Ser Leu Ile Gly Tyr Ala Ser Leu Arg
 370 375 380
 Leu His Tyr Val Thr Val Lys Lys Pro Thr Ala Val Asp Pro Asn Ser
 385 390 395 400
 Ile Val Glu Cys Arg Val Gly Asp Gly Thr Val Leu Gly Thr Gly Val
 405 410 415
 Gly Arg Asn Ile Lys Ile Ala Gly Ile Arg Ala Ala Glu Asn Ala Leu
 420 425 430
 Arg Asp Lys Lys Met Leu Asp Phe Tyr Ala Lys Gln Arg Ala Ala Ile
 435 440 445
 Pro Arg Ser Glu Ser Val Leu Lys Asp Pro Ser Gln Lys Asn Lys Lys
 450 455 460
 Arg Lys Phe Ser Asp Thr Ser
 465 470

<210> 6
 <211> 226
 <212> PRT
 <213> Escherichia coli

<400> 6

Met Asn Pro Ile Val Ile Asn Arg Leu Gln Arg Lys Leu Gly Tyr Thr
 1 5 10 15
 Phe Asn His Gln Glu Leu Leu Gln Gln Ala Leu Thr His Arg Ser Ala
 20 25 30
 Ser Ser Lys His Asn Glu Arg Leu Glu Phe Leu Gly Asp Ser Ile Leu
 35 40 45
 Ser Tyr Val Ile Ala Asn Ala Leu Tyr His Arg Phe Pro Arg Val Asp
 50 55 60
 Glu Gly Asp Met Ser Arg Met Arg Ala Thr Leu Val Arg Gly Asn Thr
 65 70 75 80

Leu Ala Glu Leu Ala Arg Glu Phe Glu Leu Gly Glu Cys Leu Arg Leu
 85 90 95
 Gly Pro Gly Glu Leu Lys Ser Gly Gly Phe Arg Arg Glu Ser Ile Leu
 100 105 110
 Ala Asp Thr Val Glu Ala Leu Ile Gly Gly Val Phe Leu Asp Ser Asp
 115 120 125
 Ile Gln Thr Val Glu Lys Leu Ile Leu Asn Trp Tyr Gln Thr Arg Leu
 130 135 140
 Asp Glu Ile Ser Pro Gly Asp Lys Gln Lys Asp Pro Lys Thr Arg Leu
 145 150 155 160
 Gln Glu Tyr Leu Gln Gly Arg His Leu Pro Leu Pro Thr Tyr Leu Val
 165 170 175
 Val Gln Val Arg Gly Glu Ala His Asp Gln Glu Phe Thr Ile His Cys
 180 185 190
 Gln Val Ser Gly Leu Ser Glu Pro Val Val Gly Thr Gly Ser Ser Arg
 195 200 205
 Arg Lys Ala Glu Gln Ala Ala Ala Glu Gln Ala Leu Lys Lys Leu Glu
 210 215 220
 Leu Glu
 225

<210> 7
 <211> 11
 <212> PRT
 <213> Homo sapiens

<400> 7

His Asn Glu Arg Leu Glu Phe Leu Gly Asp Ser
 1 5 10

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 <213> Artificial
 <220>
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20

<210> 9
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<220>
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<400> 9
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<210> 10
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<400> 10
 cggatcatta aagagcaagc 20

<210> 11
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<400> 11
 tattcaccaa agagcttcgc 20

<210> 12
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<400> 12
 caatcgtgga aagaagcaga 20

<210> 13
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<400> 13
 gctcccattt ccgcttgctg 20

<210> 14
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<400> 14	
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tgcacattca ccaaagtcaa	20
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agtctagggt cacaatctgg	20
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<210> 20
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<400> 20
cattaattct cgcagctagc gctgcgttct tcatcgacgc 40

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<400> 21
ccaaatactg atcgacaact tattgaaact tctcc 35

<210> 22
<211> 37
<212> DNA
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<220>
<223> Synthetic

<400> 22
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<210> 23
<211> 27
<212> DNA
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<220>
<223> Synthetic

<400> 23
tcgacttctg gcaagggcat tcacatt 27

<210> 24
<211> 26
<212> DNA
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<400> 24
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<210> 25
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 <400> 25
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 <400> 26
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 <210> 27
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 <400> 27
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 <210> 28
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 <400> 28
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 <210> 29
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 <400> 29
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<211> 26
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 <400> 30
 cactgggcag gaaagaacta gggttg 26

 <210> 31
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 <400> 31
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 <210> 32
 <211> 50
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 <400> 32
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 <210> 33
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 <212> DNA
 <213> Artificial
 <220>
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 <400> 33
 caaggcacgc ctctcagatc gctagagaag gcttttctca 40

 <210> 34
 <211> 40
 <212> DNA
 <213> Artificial
 <220>
 <223> Synthetic

 <400> 34
 cattaattct cgcagctagc gctgcgttct tcatcgacgc 40

 <210> 35
 <211> 20
 <212> PRT

<213> Homo sapiens

<400> 35

Cys Arg Ser Asp Tyr Asp Arg Gly Arg Thr Pro Ser Arg His Arg Ser
1 5 10 15

Tyr Glu Arg Ser
20

<210> 36

<211> 20

<212> PRT

<213> Homo sapiens

<400> 36

Cys Arg Trp Glu Arg Glu His Gln Glu Arg Glu Pro Asp Glu Thr Glu
1 5 10 15

Asp Ile Lys Lys
20

19

19